SEQUENCE LISTING

<110>	Immunokick Inc.	J	C20 Rec'd P	CT/PTO	o 5 MA	200 5
<120>	TRANSGENIC MAMMAL CARRYING OUTILIZATION THEREOF	GANP GENE	TRANSFERRED	THEREI	NTO AND	
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	PCT/JP02/11598 2002-11-07					
<160>	36					
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	t ttc r Phe												_	_		701
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110	115	120

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	140	_,_		-,-		145					150	-,-				
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σσσ	cct	gag	tca	gag	cca	gaa	222	acc	cao	age	can	att	tet	tet	gga	893
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155	110	Olu	561	oru	160	oru	Lys	1111	OIII	165	OIII	116	361	261		
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+++	+++			+	+						***					0.41
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Met		Pro	Ser	Arg	Leu		Ala	Val	Thr	Lys		Glu	Glu	Glu	Ser	
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oru	Lcu	1111	430	110	0111	Oy 3	Lys	435	110		пор	.,.	440	71511	пор	
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540

550

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680

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Lys Glu Tyr Ser Arg Ser Ser Ala Asp Gln Glu Glu Pro Leu Pro His

675

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830	835	840

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			cgg Arg			 _	_				3197
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			gga Gly	 _	_		_		Arg		3293
			aac Asn							_	3341

ctg	gct	acg g	gag o	ctg	ccc a	atc a	agc ac	ct ca	ag ag	ga go	ct g	gt gga	a ga	ас сса	3389
Leu	Ala	Thr (Glu I	Leu l	Pro]	lle S	Ser Th	nr Gl	ln Aı	g Al	la G	ly Gly	7 As	sp Pro)
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1110	1115	1120

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	_	gag Glu 1215							gaa Glu	4064
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Pro	${\tt Pro}$	Lys	Val	Lys	Ser	Glu	Glu	Val	Ala	Glu	Glu	Glu	Leu	Ser	
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	cta Leu									_		5369

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		ttt Phe									5729

5774

1775

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1785	1790	1795
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1920

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Leu Ala Phe Ser Gln Val Pro Ser Phe Ala Thr Pro Ser Gly Gly Ser His Ser Ser Ser Leu Pro Ala Phe Gly Leu Thr Gln Thr Ser Ser Val Gly Leu Phe Ser Ser Leu Glu Ser Thr Pro Ser Phe Ala Ala Thr Ser Ser Ser Ser Val Pro Gly Asn Thr Ala Phe Ser Phe Lys Ser Thr Ser Ser Val Gly Val Phe Pro Ser Gly Ala Thr Phe Gly Pro Glu Thr Gly Glu Val Ala Gly Ser Gly Phe Arg Lys Thr Glu Phe Lys Phe Lys Pro Leu Glu Asn Ala Val Phe Lys Pro Ile Pro Gly Pro Glu Ser Glu Pro Glu Lys Thr Gln Ser Gln Ile Ser Ser Gly Phe Phe Thr Phe Ser His Pro Val Gly Ser Gly Ser Gly Gly Leu Thr Pro Phe Ser Phe Pro Gln

Val Thr Asn Ser Ser Val Thr Ser Ser Ser Phe Ile Phe Ser Lys Pro Val Thr Ser Asn Thr Pro Ala Phe Ala Ser Pro Leu Ser Asn Gln Asn Val Glu Glu Glu Lys Arg Val Ser Thr Ser Ala Phe Gly Ser Ser Asn Ser Ser Phe Ser Thr Phe Pro Thr Ala Ser Pro Gly Ser Leu Gly Glu Pro Phe Pro Ala Asn Lys Pro Ser Leu Arg Gln Gly Cys Glu Glu Ala Ile Ser Gln Val Glu Pro Leu Pro Thr Leu Met Lys Gly Leu Lys Arg Lys Glu Asp Gln Asp Arg Ser Pro Arg Arg His Cys His Glu Ala Ala Glu Asp Pro Asp Pro Leu Ser Arg Gly Asp His Pro Pro Asp Lys Arg . 305 Pro Val Arg Leu Asn Arg Pro Arg Gly Gly Thr Leu Phe Gly Arg Thr

Ile Gln Glu Val Phe Lys Ser Asn Lys Glu Ala Gly Arg Leu Gly Ser Lys Glu Ser Lys Glu Ser Gly Phe Ala Glu Pro Gly Glu Ser Asp His Ala Ala Val Pro Gly Gly Ser Gln Ser Thr Met Val Pro Ser Arg Leu Pro Ala Val Thr Lys Glu Glu Glu Glu Ser Arg Asp Glu Lys Glu Asp Ser Leu Arg Gly Lys Ser Val Arg Gln Ser Lys Arg Arg Glu Glu Trp Ile Tyr Ser Leu Gly Gly Val Ser Ser Leu Glu Leu Thr Ala Ile Gln Cys Lys Asn Ile Pro Asp Tyr Leu Asn Asp Arg Ala Ile Leu Glu Lys His Phe Ser Lys Ile Ala Lys Val Gln Arg Val Phe Thr Arg Arg Ser

Lys Lys Leu Ala Val Ile His Phe Phe Asp His Ala Ser Ala Ala Leu

Ala Arg Lys Lys Gly Lys Gly Leu His Lys Asp Val Val Ile Phe Trp His Lys Lys Lys Ile Ser Pro Ser Lys Lys Leu Phe Pro Leu Lys Glu Lys Leu Gly Glu Ser Glu Ala Ser Gln Gly Ile Glu Asp Ser Pro Phe Gln His Ser Pro Leu Ser Lys Pro Ile Val Arg Pro Ala Ala Gly Ser Leu Leu Ser Lys Ser Ser Pro Val Lys Lys Pro Ser Leu Leu Lys Met His Gln Phe Glu Ala Asp Pro Phe Asp Ser Gly Ser Glu Gly Ser Glu Gly Leu Gly Ser Cys Val Ser Ser Leu Ser Thr Leu Ile Gly Thr Val Ala Asp Thr Ser Glu Glu Lys Tyr Arg Leu Leu Asp Gln Arg Asp Arg

Ile Met Arg Gln Ala Arg Val Lys Arg Thr Asp Leu Asp Lys Ala Arg

Ala Phe Val Gly Thr Cys Pro Asp Met Cys Pro Glu Lys Glu Arg Tyr Leu Arg Glu Thr Arg Ser Gln Leu Ser Val Phe Glu Val Val Pro Gly Thr Asp Gln Val Asp His Ala Ala Ala Val Lys Glu Tyr Ser Arg Ser Ser Ala Asp Gln Glu Glu Pro Leu Pro His Glu Leu Arg Pro Ser Ala Val Leu Ser Arg Thr Met Asp Tyr Leu Val Thr Gln Ile Met Asp Gln Lys Glu Gly Ser Leu Arg Asp Trp Tyr Asp Phe Val Trp Asn Arg Thr Arg Gly Ile Arg Lys Asp Ile Thr Gln Gln His Leu Cys Asp Pro Leu 730 · Thr Val Ser Leu Ile Glu Lys Cys Thr Arg Phe His Ile His Cys Ala

His Phe Met Cys Glu Glu Pro Met Ser Ser Phe Asp Ala Lys Ile Asn

Asp Leu Arg Asn Lys Gly Val Phe Cys Ala Ser Glu Ala Glu Phe Gln Gly Tyr Asn Val Leu Leu Asn Leu Asn Lys Gly Asp Ile Leu Arg Glu Val Gln Gln Phe His Pro Asp Val Arg Asn Ser Pro Glu Val Asn Phe Ala Val Gln Ala Phe Ala Ala Leu Asn Ser Asn Asn Phe Val Arg Phe Phe Lys Leu Val Gln Ser Ala Ser Tyr Leu Asn Ala Cys Leu Leu His Cys Tyr Phe Asn Gln Ile Arg Lys Asp Ala Leu Arg Ala Leu Asn Val Ala Tyr Thr Val Ser Thr Gln Arg Ser Thr Val Phe Pro Leu Asp Gly Val Val Arg Met Leu Leu Phe Arg Asp Ser Glu Glu Ala Thr Asn Phe

Asn Glu Asn Met Thr Lys Cys Leu Gln Ser Leu Lys Glu Met Tyr Gln

Leu Asn Tyr His Gly Leu Thr Val Ala Asp Gly Cys Val Glu Leu Asn 915 920 925

Arg Ser Ala Phe Leu Glu Pro Glu Gly Leu Cys Lys Ala Arg Lys Ser 930 935 940

Val Phe Ile Gly Arg Lys Leu Thr Val Ser Val Gly Glu Val Val Asn 945 950 955 960

Gly Gly Pro Leu Pro Pro Val Pro Arg His Thr Pro Val Cys Ser Phe.
965 970 975

Asn Ser Gln Asn Lys Tyr Val Gly Glu Ser Leu Ala Thr Glu Leu Pro 980 985 990

Ile Ser Thr Gln Arg Ala Gly Gly Asp Pro Ala Gly Gly Gly Arg Gly 995 1000 1005

Glu Asp Cys Glu Ala Glu Val Asp Leu Pro Thr Leu Ala Val Leu 1010 1015 1020

Pro Gln Pro Pro Pro Ala Ser Ser Ala Thr Pro Ala Leu His Val 1025 1030 . 1035

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- Asp Ser Asp Leu Val Gln Val Val Asp Glu Leu Ile Gln Glu Ala 1070 1075 1080
- Leu Gln Val Asp Cys Glu Glu Val Ser Ser Ala Gly Ala Ala Tyr 1085 1090 1095
- Val Ala Ala Ala Leu Gly Val Ser Asn Ala Ala Val Glu Asp Leu 1100 1105 1110
- Ile Thr Ala Ala Thr Thr Gly Ile Leu Arg His Val Ala Ala Glu 1115 1120 1125
- Glu Val Ser Met Glu Arg Gln Arg Leu Glu Glu Glu Lys Gln Arg 1130 1135 1140
- Ala Glu Glu Glu Arg Leu Lys Gln Glu Arg Glu Leu Met Leu Thr 1145 1150 1155
- Gln Leu Ser Glu Gly Leu Ala Ala Glu Leu Thr Glu Leu Thr Val 1160 1165 1170
- Thr Glu Cys Val Trp Glu Thr Cys Ser Gln Glu Leu Gln Ser Ala 1175 1180 1185

- Val Lys Ile Asp Gln Lys Val Arg Val Ala Arg Cys Cys Glu Ala 1190 1195 1200
- Val Cys Ala His Leu Val Asp Leu Phe Leu Ala Glu Glu Ile Phe 1205 1210 1215
- Gln Thr Ala Lys Glu Thr Leu Gln Glu Leu Gln Cys Phe Cys Lys 1220 1225 . 1230
- Tyr Leu Gln Arg Trp Arg Glu Ala Val Ala Ala Arg Lys Lys Phe 1235 1240 1245
- Arg Arg Gln Met Arg Ala Phe Pro Ala Ala Pro Cys Cys Val Asp 1250 1255 1260
- Val Asn Asp Arg Leu Gln Ala Leu Val Pro Ser Ala Glu Cys Pro 1265 1270 1275
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- Ala Gly Lys Val Gly Val Ser Cys Thr Arg Leu Arg Arg Leu Arg 1295 1300 1305
- Asn Lys Thr Ala His Gln Ile Lys Val Gln His Phe His Gln Gln 1310 1315 1320

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- Asp Asp Ser Met Val Gly Asp Ile Gly Asp Asn Ala Gly Asp Ile
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- Thr Val Ser Val Asn Val Cys Ile Lys Val Ala His Gly Thr Leu 1415 1420 1425
- Ser Asp Ser Ala Leu Asp Ala Val Glu Thr Gln Lys Asp Leu Leu 1430 1435 1440
- Gly Thr Ser Gly Leu Met Leu Leu Leu Pro Pro Lys Val Lys Ser 1445 1450 1455

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- Glu Glu Val Ala Glu Glu Glu Leu Ser Trp Leu Ser Ala Leu Leu 1460 1465 1470
- Gln Leu Lys Gln Leu Leu Gln Ala Lys Pro Phe Gln Pro Ala Leu 1475 1480 1485
- Pro Leu Val Val Leu Val Pro Ser Ser Arg Gly Asp Ser Ala Gly 1490 1495 1500
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- Lys Leu Ile Ser Asp Tyr Ile Val Val Glu Ile Pro Asp Ser Val 1520 1525 1530
- Asn Asp Leu Gln Gly Thr Val Lys Val Ser Gly Ala Val Gln Trp 1535 1540 1545
- Leu Ile Ser Gly Cys Pro Gln Ala Leu Asp Leu Cys Cys Gln Thr 1550 1555 1560
- Leu Val Gln Tyr Val Glu Asp Gly Ile Ser Arg Glu Phe Ser Arg 1565 1570 1575
- Arg Phe Phe His Asp Arg Arg Glu Arg Arg Leu Ala Ser Leu Pro 1580 1585 1590

- Ser Gln Glu Pro Ser Thr Ile Ile Glu Leu Phe Asn Ser Val Leu 1595 1600 1605
- Gln Phe Leu Ala Ser Val Val Ser Ser Glu Gln Leu Cys Asp Ile 1610 1615 1620
- Ser Trp Pro Val Met Glu Phe Ala Glu Val Gly Gly Ser Gln Leu 1625 1630 1635
- Leu Pro His Leu His Trp Asn Ser Pro Glu His Leu Ala Trp Leu 1640 1645 1650
- Lys Gln Ala Val Leu Gly Phe Gln Leu Pro Gln Met Asp Leu Pro 1655 1660 1665
- Pro Pro Gly Ala Pro Trp Leu Pro Val Cys Ser Met Val Ile Gln 1670 1675 1680
- Tyr Thr Ser Gln Ile Pro Ser Ser Ser Gln Thr Gln Pro Val Leu 1685 1690 1695
- Gln Ser Gln Ala Glu Asn Leu Leu Cys Arg Thr Tyr Gln Lys Trp 1700 1705 1710
- Lys Asn Lys Ser Leu Ser Pro Gly Gln Glu Leu Gly Pro Ser Val 1715 1720 1725

- Ala Glu Ile Pro Trp Asp Asp Ile Ile Thr Leu Cys Ile Asn His 1730 1735 1740
- Lys Leu Arg Asp Trp Thr Pro Pro Arg Leu Pro Val Thr Leu Glu 1745 1750 1755
- Ala Leu Ser Glu Asp Gly Gln Ile Cys Val Tyr Phe Phe Lys Asn 1760 1765 1770
- Leu Leu Arg Lys Tyr His Val Pro Ser Ser Trp Glu Gln Ala Arg 1775 1780 1785
- Met Gln Thr Gln Arg Glu Leu Gln Leu Ser His Gly Arg Ser Gly 1790 1795 1800
- Met Arg Ser Ile His Pro Pro Thr Ser Thr Phe Pro Thr Pro Leu 1805 1810 1815
- Leu His Val His Gln Lys Gly Lys Lys Glu Glu Ser Gly Arg 1820 1825 1830
- Glu Gly Ser Leu Ser Thr Glu Asp Leu Leu Arg Gly Ala Ser Ala 1835 1840 1845
- Glu Glu Leu Leu Ala Gln Ser Leu Ser Ser Ser Leu Leu Glu Glu 1850 1855 1860

Lys Glu Glu Asn Lys Arg Phe Glu Asp Gln Leu Gln Gln Trp Leu 1865 1870 1875

Ser Gln Asp Ser Gln Ala Phe Thr Glu Ser Thr Arg Leu Pro Leu 1880 1885 1890

Tyr Leu Pro Gln Thr Leu Val Ser Phe Pro Asp Ser Ile Lys Thr 1895 1900 1905

Gln Thr Met Val Lys Thr Ser Thr Ser Pro Gln Asn Ser Gly Thr 1910 1915 1920

Gly Lys Gln Leu Arg Phe Ser Glu Ala Ser Gly Ser Ser Leu Thr 1925 1930 1935

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Glu Glu Ala Ala Ser Glu Leu His Leu Ser Ala Leu Leu Glu Met 1955 1960 1965

Val Asp Met 1970

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									4		+ 0 +	+ 0 +	o a t	aat	ata	103
ttc a																100
Phe S	ser	Gly	10	GIN	Pro	Ser	нта	15	Sei	міа	361	561	20	11311	101	
			10					10								
gga a	ara	ctt	cca	tet	aag	CCE	cca	ttt	cga	ttt	ggt	caa	cct	tct	ctt	151
Gly 1																
01,		25			_,		30		_			35				
ttt į	gga	caa	aac	agt	acc	tta	tct	ggg	aag	agc	tcg	gga	ttt	tca	cag	199
Phe (Gly	Gln	Asn	Ser	Thr	Leu	Ser	Gly	Ĺуѕ	Ser	Ser	Gly	Phe	Ser	Gln	
4	40					45					50					
gta																247
Val :	Ser	Ser	Phe	Pro	Ala	Ser	Ser	Gly	Val	Ser	His	Ser	Ser	Ser		
55					60					65					70	
																205
caa																295
Gln	Thr	Leu	Gly		Thr	Gln	Thr	Ser		Val	Gly	Pro	Pne		GIY	
				75					80					85		
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ctt																040
Leu	GIU	пls		ser	ınr	rne	val	95	1111	261	019	110	100		501	
			90					55					_ 0			

												CCC				391
Ser	Val	Leu 105	Gly	Asn	lhr	Gly	110	Ser	Pne	Lys	Ser	Pro 115	Inr	Ser	vaı	
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Gly	Ala	Phe	Pro	Ser	Thr	Ser	Ala	Phe	Gly	Gln	Glu	Ala	Gly	Glu	Ile	
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Val	Asn	Ser	Gly	Phe	Gly	Lys	Thr	Glu	Phe	Ser	Phe	Lys	Pro	Leu	Glu	
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Asn	Ala	Val	Phe	Lys	Pro	Ile	Leu	Gly	Ala	Glu	Ser	Glu	Pro	Glu	Lys	
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Thr	Gln	Ser	Gln	Ile	Ala	Ser	Gly	Phe	Phe	Thr	Phe	Ser	His	Pro	Ile	
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Ser	Ser		Pro	Gly	GIY	Leu	190	Pro	Pne	Ser	rne	Pro 195	GIN	vai	inr	
		185					190					190				
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Ser	Ser	Ser	Ala	Thr	Thr	Ser	Asn	Phe	Thr	Phe	Ser	Lys	Pro	Val	Ser	
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agt	aat	aat	tca	tta	tct	gcc	ttt	acc	cct	gct	ttg	tca	aac	caa	aat	727
Ser	Asn	Asn	Ser	Leu	Ser	Ala	Phe	Thr	Pro	Ala	Leu	Ser	Asn	Gln	Asn	
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								gaa Glu		871
								aaa Lys		919
_								cca Pro		967
								aaa Lys		1015
_	_	_						cgg Arg 340		1063
•	•	_		_				ggc Gly		1111
								agt Ser		1159
								tcc Ser		1207

				gaa Glu					1255
				ccg Pro					1303
				ttg Leu					1351
_	_			tac Tyr 445					1399
				aaa Lys					1447
				cat His					1495
				agt Ser					1543
				ccc Pro					 1591
				gaa Glu 525					1639

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			gga Gly		•				1	783
			tgt Cys						1	831
			aag Lys						1	879
			gct Ala 620						. 1	927
			acc Thr						1	975
			cgt Arg							2023
			gac Asp					_	2	2071

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Ser		Ala	Asp	GIn	Glu		Pro	Leu	Pro	HIS		Leu	Arg	Pro	Leu	
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Pro	Val	Leu	Ser	Arg	Thr	Met	Asp	Tyr	Leu	Val	Thr	Gln	Ile	Met	Asp	
695					700					705					710	
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cag	aag	gag	ggc	agc	ctg	cgg	gat	tgg	tat	gac	ttc	gtg	tgg	aac	cgc	2215
					Leu											
	•		•	715		_	_	_	720					725		
acσ	cat	ggc	ata	Cgg	aag	gat.	atc	acg	cag	cag	cac	ctc	tgt	gac	ccc	2263
					Lys											
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Leu	Ihr		Ser	Leu	Ile	GIU		Cys	ınr	Arg	rne		116	піѕ	Cys	
		745					750					755		4		
																0250
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Ala	His	Phe	Met	Cys	Glu		Pro	Met	Ser	Ser		Asp	Ala	Lys	lle	
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Gln	Asp	Leu	Arg	Asn	Lys	Gly	Val	Phe	Cys	Ala	Ser	Glu	Ala	Glu	Phe	
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		ttt Phe						2599
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		cag Gln						2695
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		ctg Leu						2791
		ggc Gly						2839
		ctg Leu 940						2887
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	cca Pro 1030	G1n			gcc Ala 1035							3166
	cct Pro 1045	Val			acc Thr 1050					-		3211
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		Sei			ctg Leu 1080	Ala						3301
		Ala			gac Asp 1095	Cys			ggc Gly 1100			3346

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Arg	Val	Met	Met	Glu	Phe	Val		Glu	Thr	Cys			Glu	Leu	
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_	aat	•				_									3661
Lys	Asn	Ala	Val	Glu	Ihr			Arg	vai	Arg			Arg	Cys	
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Cys	Glu	Asp	vai	Cys	Ala			vai	ASP	Leu	1220		Val	Olu	
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_						aag									0.01
Glu	Ile	rne	GIN	ınr	чта			1111	Leu	9111	1235		0111	Cys	
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						atg Met 1260					3841
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		Leu				cac His 1350					4111
		Lys				gtg. Val 1365			Glu		4156

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Asp	Leu	Pro	Pro	Leu	Gly	Ala	Pro	Trp	Leu	Pro	Val	Cys	Ser	Met	
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Val	Val	Gln	Tyr	Ala	Ser	Gln	Ile	Pro	Ser	Ser	Arg	Gln	Thr	Gln	
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Pro	Val	Leu	Gln	Ser	Gln	Val	Glu	Asn	Leu	Leu	His	Arg	Thr	Tyr	
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Cys	Arg	Trp	Lys	Ser	Lys	Ser	Pro	Ser	Pro	Val	His	Gly	Ala	Gly	
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Pro	Ser	Val	Met	Glu	Ile	Pro	Trp	Asp	Asp	Leu	Ile	Ala	Leu	Cys	
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acg tg	t cta	ggc	gaa	cga	cta	aag	cac	ctg	gaa	agg	ctg	atc	cgg	5911
Thr Cy	s Leu	Gly	${\tt Glu}$	Arg	Leu	Lys	His	Leu	Glu	Arg	Leu	Ile	Arg	
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Ser Se	r Arg	Glu	Glu	Glu	Val	Ala	Ser	Glu	Leu	His	Leu	Ser	Ala	
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Leu Le	u Asp	Met	Val	Asp	Ile									
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Ser Phe Lys Pro Leu Glu Asn Ala Val Phe Lys Pro Ile Leu Gly Ala

Glu Ser Glu Pro Glu Lys Thr Gln Ser Gln Ile Ala Ser Gly Phe Phe 165 170 175

Thr Phe Ser His Pro Ile Ser Ser Ala Pro Gly Gly Leu Ala Pro Phe 180 185 190

Ser Phe Pro Gln Val Thr Ser Ser Ser Ala Thr Thr Ser Asn Phe Thr 195 200 205

Phe Ser Lys Pro Val Ser Ser Asn Asn Ser Leu Ser Ala Phe Thr Pro 210 215 220

Ala Leu Ser Asn Gln Asn Val Glu Glu Glu Lys Arg Gly Pro Lys Ser 225 230 235 240

Ile Phe Gly Ser Ser Asn Asn Ser Phe Ser Ser Phe Pro Val Ser Ser 245 250 255

Ala Val Leu Gly Glu Pro Phe Gln Ala Ser Lys Ala Gly Val Arg Gln 260 265 270

Gly Cys Glu Glu Ala Val Ser Gln Val Glu Pro Leu Pro Ser Leu Met 275 280 285

Lys Gly Leu Lys Arg Lys Glu Asp Gln Asp Arg Ser Pro Arg Arg His 290 295 300 Gly His Glu Pro Ala Glu Asp Ser Asp Pro Leu Ser Arg Gly Asp His 305 310 315 320

Pro Pro Asp Lys Arg Pro Val Arg Leu Asn Arg Pro Arg Gly Gly Thr 325 330 335

Leu Phe Gly Arg Thr Ile Gln Asp Val Phe Lys Ser Asn Lys Glu Val 340 345 350

Gly Arg Leu Gly Asn Lys Glu Ala Lys Lys Glu Thr Gly Phe Val Glu 355 360 365

Ser Ala Glu Ser Asp His Met Ala Ile Pro Gly Gly Asn Gln Ser Val 370 375 380

Leu Ala Pro Ser Arg Ile Pro Gly Val Asn Lys Glu Glu Glu Thr Glu 385 390 395 400

Ser Arg Glu Lys Lys Glu Asp Ser Leu Arg Gly Thr Pro Ala Arg Gln
405 410 415

Ser Asn Arg Ser Glu Ser Thr Asp Ser Leu Gly Gly Leu Ser Pro Ser 420 425 430

Glu Val Thr Ala Ile Gln Cys Lys Asn Ile Pro Asp Tyr Leu Asn Asp 435 440 445 Arg Thr Ile Leu Glu Asn His Phe Gly Lys Ile Ala Lys Val Gln Arg Ile Phe Thr Arg Arg Ser Lys Lys Leu Ala Val His Phe Phe Asp His Ala Ser Ala Ala Leu Ala Arg Lys Lys Gly Lys Ser Leu His Lys Asp Met Ala Ile Phe Trp His Arg Lys Lys Ile Ser Pro Asn Lys Lys Pro Phe Ser Leu Lys Glu Lys Lys Pro Gly Asp Gly Glu Val Ser Pro Ser Thr Glu Asp Ala Pro Phe Gln His Ser Pro Leu Gly Lys Ala Ala Gly Arg Thr Gly Ala Ser Ser Leu Leu Asn Lys Ser Ser Pro Val Lys Lys Pro Ser Leu Leu Lys Ala His Gln Phe Glu Gly Asp Ser Phe Asp

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Ser Thr Leu Ile Gly Thr Val Ala Glu Thr Ser Lys Glu Lys Tyr Arg 595 600 605

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Thr Asp Leu Asp Lys Ala Arg Thr Phe Val Gly Thr Cys Leu Asp Met 625 630 635 640

Cys Pro Glu Lys Glu Arg Tyr Met Arg Glu Thr Arg Ser Gln Leu Ser 645 650 655

Val Phe Glu Val Val Pro Gly Thr Asp Gln Val Asp His Ala Ala Ala 660 665 670

Val Lys Glu Tyr Ser Arg Ser Ser Ala Asp Gln Glu Glu Pro Leu Pro 675 680 685

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Gln His Leu Cys Asp Pro Leu Thr Val Ser Leu Ile Glu Lys Cys Thr 740 745 750

Arg Phe His Ile His Cys Ala His Phe Met Cys Glu Glu Pro Met Ser 755 760 765

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- Cys Leu Ser Ser Ser Leu Leu Glu Lys Glu Glu Asn Lys Arg 1865 1870 1875
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- Phe Thr Asp Leu Thr Ser Leu Pro Leu Tyr Leu Pro Gln Thr Leu 1895 1900 1905
- Val Ser Leu Ser His Thr Ile Glu Pro Val Met Lys Thr Ser Val 1910 1915 1920
- Thr Thr Ser Pro Gln Ser Asp Met Met Arg Glu Gln Leu Gln Leu 1925 1930 1935
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